# **Resource Summary Report**

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# **LCRC Proteomics Core Facility**

RRID:SCR\_009894

Type: Tool

### **Proper Citation**

LCRC Proteomics Core Facility (RRID:SCR\_009894)

#### **Resource Information**

**URL:** http://xula.eagle-i.net/i/00000135-81af-5452-a9f8-d64580000000

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**Description:** Core facility that provides the following services: Mass spectrometry services. The mission of Proteomics Core Facility is to help researchers advance their biomedical research programs. The Facility is equipped with gel electrophoresis units, image analyzers and digitizers, and robotics for mass spectrometry sample preparation. Currently, a Matrix-Assisted Laser Desorption/Ionization Mass Spectrometer instrument implements protein identification. The various applications include the studies of protein expression profile, posttranslational modifications, and partial sequencing of novel proteins. The staff also consult with researchers about their particular research interests and assist with the development of specific applications and solution to problems.

Resource Type: access service resource, core facility, service resource

**Keywords:** mass spectrometry assay

**Funding:** 

Resource Name: LCRC Proteomics Core Facility

Resource ID: SCR\_009894

Alternate IDs: nlx\_156374

**Record Creation Time:** 20220129T080255+0000

Record Last Update: 20250407T215833+0000

## Ratings and Alerts

No rating or validation information has been found for LCRC Proteomics Core Facility.

No alerts have been found for LCRC Proteomics Core Facility.

### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We have not found any literature mentions for this resource.